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**IN THE UNITED STATES PATENT AND TRADEMARK OFFICE**

**Atty. Docket No: 37629-0004**

*In re* patent application of  
KNAPPIK, Achim *et al.*

Application No.: 09/490,324

Filed: January 24, 2000

For: PROTEIN/ (POLY) PEPTIDE LIBRARIES



Group Art Unit: 1631

Examiner: M. Zeman

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Plunkett

4/10/02

**AMENDMENT ACCOMPANYING FORMAL DRAWINGS**

Commissioner for Patents  
Washington, D.C. 20231

Sir:

Applicants respectfully request that the following amendments be entered into the application:

**IN THE SPECIFICATION:**

**Page 18, line 4, paragraph 2:** Please amend as follows:

*B1* *SUS* *CI* Figs. 2A-2G: Alignment of consensus sequences designed for each subgroup (amino acid residues are shown with their standard one-letter abbreviation). (2A-2B) kappa sequences, (2C-2D) lambda sequences and (2E-2G), heavy chain sequences. The positions are numbered according to Kabat (1991). In order to maximize homology in the alignment, gaps (-) have been introduced in the sequence at certain positions.

**Page 18, line 10, paragraph 3:** Please amend as follows:

Figs. 3A-3K: Gene sequences of the synthetic V kappa consensus genes. The corresponding amino acid sequences (see Figs. 2A-2B) as well as the unique cleavage sites are also shown.

**Page 18, line 13, paragraph 4:** Please amend as follows:

Figs. 4A-4I: Gene sequences of the synthetic V lambda consensus genes. The corresponding amino acid sequences (see Figs. 2C-2D) as well as the unique cleavage sites are also shown.

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